RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _10/677,983 A_

Source: _1 FWD_

Date Processed by STIC: _1/6/05_

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RAW SEQUENCE LISTING   DATE: 01/06/2005
PATENT APPLICATION: US/10/677,983A   TIME: 15:57:28

Input Set : A:\Felder39.app
Output Set: N:\CRF4\01062005\J677983A.raw

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 3 <110> APPLICANT: FELDER, ROBIN A.
 4         JOSE, PEDRO
 6 <120> TITLE OF INVENTION: G PROTEIN-RELATED KINASE MUTANTS IN ESSENTIAL
 7         HYPERTENSION
 9 <130> FILE REFERENCE: FELDER 3.9-001 CONT DIV
11 <140> CURRENT APPLICATION NUMBER: 10/677,983A
12 <141> CURRENT FILING DATE: 2003-10-02
14 <150> PRIOR APPLICATION NUMBER: 09/614,748
15 <151> PRIOR FILING DATE: 2000-07-12
17 <150> PRIOR APPLICATION NUMBER: PCT/US99/00663
18 <151> PRIOR FILING DATE: 1999-01-12
20 <150> PRIOR APPLICATION NUMBER: 60/071,199
21 <151> PRIOR FILING DATE: 1998-01-12
23 <150> PRIOR APPLICATION NUMBER: 60/098,279
24 <151> PRIOR FILING DATE: 1998-08-28
26 <160> NUMBER OF SEQ ID NOS: 34
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45 Ile Glu Lys Asp Tyr Ser Ser Leu Cys Asp Lys Gln Pro Ile Gly Arg
46      50                  55                  60
48 Arg Leu Phe Arg Gln Phe Cys Asp Thr Lys Pro Thr Leu Lys Arg His
49 65                  70                  75                  80
51 Ile Glu Phe Leu Asp Ala Val Ala Glu Tyr Glu Val Ala Asp Asp Glu
52              85                  90                  95
54 Asp Arg Ser Asp Cys Gly Leu Ser Ile Leu Asp Arg Phe Phe Asn Asp
55          100                 105                 110
57 Lys Leu Ala Ala Pro Leu Pro Glu Ile Pro Pro Asp Val Val Thr Glu
58          115                 120                 125
60 Cys Arg Leu Gly Leu Lys Glu Glu Asn Pro Ser Lys Lys Ala Phe Glu
61          130                 135                 140
63 Glu Cys Thr Arg Val Ala His Asn Tyr Leu Arg Gly Glu Pro Phe Glu
64 145                 150                 155                 160
66 Glu Tyr Gln Glu Ser Ser Tyr Phe Ser Gln Phe Leu Gln Trp Lys Trp
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73           195               200               205
75 Thr Gly Lys Met Tyr Ala Cys Lys Lys Leu Gln Lys Lys Arg Ile Lys
76      210               215               220
78 Lys Arg Lys Gly Glu Ala Met Ala Leu Asn Glu Lys Arg Ile Leu Glu
79 225               230               235               240
81 Lys Val Gln Ser Arg Phe Val Val Ser Leu Ala Tyr Ala Tyr Glu Thr
82           245               250               255
84 Lys Asp Ala Leu Cys Leu Val Leu Thr Ile Met Asn Gly Gly Asp Leu
85           260               265               270
87 Lys Phe His Ile Tyr Asn Leu Gly Asn Pro Gly Phe Asp Glu Gln Arg
88           275               280               285
90 Ala Val Phe Tyr Ala Ala Glu Leu Cys Cys Gly Leu Glu Asp Leu Gln
91      290               295               300
93 Arg Glu Arg Ile Val Tyr Arg Asp Leu Lys Pro Glu Asn Ile Leu Leu
94 305               310               315               320
96 Asp Asp Arg Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Thr Glu
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106      370               375               380
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112           405               410               415
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123 Pro Phe Cys Pro Asp Pro His Ala Val Tyr Cys Lys Asp Val Leu Asp
124 465               470               475               480
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133      515               520               525
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175 Glu Cys Thr Arg Val Ala His Asn Tyr Leu Arg Gly Glu Pro Phe Glu
176       115             120             125
178 Glu Tyr Gln Glu Ser Ser Tyr Phe Ser Gln Phe Leu Gln Trp Lys Trp
179     130             135             140
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182 145             150             155             160
184 Leu Gly Lys Gly Gly Phe Gly Glu Val Cys Ala Cys Gln Val Arg Ala
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187 Thr Gly Lys Met Tyr Ala Cys Lys Lys Leu Gln Lys Lys Arg Ile Lys
188         180             185             190
190 Lys Arg Lys Gly Glu Ala Met Ala Leu Asn Glu Lys Arg Ile Leu Glu
191     195             200             205
193 Lys Val Gln Ser Arg Phe Val Val Ser Leu Ala Tyr Ala Tyr Glu Thr
194     210             215             220
196 Lys Asp Ala Leu Cys Leu Val Leu Thr Ile Met Asn Gly Gly Asp Leu
197 225             230             235             240
199 Lys Phe His Ile Tyr Asn Leu Gly Asn Pro Gly Phe Asp Glu Gln Arg
200             245             250             255
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203         260             265             270
205 Arg Glu Arg Ile Val Tyr Arg Asp Leu Lys Pro Glu Asn Ile Leu Leu
206     275             280             285
208 Asp Asp Arg Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Thr Glu
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212 305             310             315             320
214 Met Ala Pro Glu Val Val Asn Asn Glu Lys Tyr Thr Phe Ser Pro Asp
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226 Ala Lys Ser Ile Cys Arg Met Leu Leu Thr Lys Asn Pro Ser Lys Arg
227 385                   390                   395                   400
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232 Phe Lys Asp Ile Asn Phe Arg Arg Leu Glu Ala Asn Met Leu Glu Pro
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248                   500                   505                   510
250 Pro Val Ser Arg Pro Asn Arg Gly Phe Phe Tyr Arg Leu Phe Arg Arg
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279 65                   70                   75                   80
281 Ile Glu Phe Leu Asp Ala Val Ala Glu Tyr Glu Val Ala Asp Asp Glu
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285             100                   105                   110
287 Lys Leu Ala Ala Pro Leu Pro Glu Ile Pro Pro Asp Val Val Thr Glu
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288           115                  120                  125
290 Cys Arg Leu Gly Leu Lys Glu Glü Asn Pro Ser Lys Lys Ala Phe Glu
291     130                  135                  140
293 Glu Cys Thr Arg Val Ala His Asn Tyr Leu Arg Gly Glu Pro Phe Glu
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305 Thr Gly Lys Met Tyr Ala Cys Lys Lys Leu Gln Lys Lys Arg Ile Lys
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326 Asp Asp Arg Gly His Ile Arg Ile Ser Asp Leu Gly Leu Ala Thr Glu
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354 465                  470                  475                  480
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VERIFICATION SUMMARY   DATE: 01/06/2005
PATENT APPLICATION: **US/10/677,983A**   TIME: 15:57:29

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